

1600

RAW SEQUENCE LISTING DATE: 05/13/2003 PATENT APPLICATION: US/09/472,067A TIME: 16:08:27

Input Set : A:\PTO.VSK.txt

Output Set: N:\CRF4\05132003\I472067A.raw

SEQUENCE LISTING

```
(1) GENERAL INFORMATION:
             (i) APPLICANT: HU, A-LI W.
                             HARTLEY, JAMES L.
                             JORDAN, HEATHER J.
      8
     10
            (ii) TITLE OF INVENTION: NUCLEIC ACID LADDERS
     12
           (iii) NUMBER OF SEQUENCES: 7
            (iv) CORRESPONDENCE ADDRESS:
     14
                   (A) ADDRESSEE: STERNE, KESSLER, GOLDSTEIN & FOX, P.L.L.C.
     15
                   (B) STREET: 1100 NEW YORK AVENUE N.W., SUITE 600
     16
     17
                   (C) CITY: WASHINGTON
                   (D) STATE: DC
     18
     19
                   (E) COUNTRY: USA
                                                                         Does Not Comply
     20
                   (F) ZIP: 20005
                                                                     Corrected Diskette Needed
             (v) COMPUTER READABLE FORM:
     22
                   (A) MEDIUM TYPE: Floppy disk
     23
                   (B) COMPUTER: IBM PC compatible
     24
                   (C) OPERATING SYSTEM: PC-DOS/MS-DOS
     25
                   (D) SOFTWARE: PatentIn Release #1.0, Version #1.30
     26
            (vi) CURRENT APPLICATION DATA:
     28
                   (A) APPLICATION NUMBER: US/09/472,067A
C-->29
                   (B) FILING DATE: 23-Dec-1999
C-->30
                   (C) CLASSIFICATION:
W --> 41
           (vii) PRIOR APPLICATION DATA:
     38
                   (A) APPLICATION NUMBER: US 09/048,242
     34
     35
                   (B) FILING DATE: 26-MAR-1998
                   (A) APPLICATION NUMBER: US 60/040,914
     39
                   (B) FILING DATE: 27-MAR-1997
     40
            (ix) TELECOMMUNICATION INFORMATION:
     43
                   (A) TELEPHONE: (202)371-2600
     44
     45
                   (B) TELEFAX: (202)371-2540
     48 (2) INFORMATION FOR SEQ ID NO: 1:
             (i) SEQUENCE CHARACTERISTICS:
                   (A) LENGTH: 10 base pairs
     51
     52
                   (B) TYPE: nucleic acid
                   (C) STRANDEDNESS: both
     53
     54
                   (D) TOPOLOGY: linear
            (ii) MOLECULE TYPE: cDNA
     56
            (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:
                                                                                   10
     64 ATCTCAGGAT
     66 (2) INFORMATION FOR SEQ ID NO: 2:
             (i) SEQUENCE CHARACTERISTICS:
     68
                   (A) LENGTH: 10 base pairs
     69
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```
(B) TYPE: nucleic acid
70
             (C) STRANDEDNESS: both
71
             (D) TOPOLOGY: linear
72
       (ii) MOLECULE TYPE: cDNA
74
       (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:
79
                                                                           10
81 ATCAGTCGAT
83 (2) INFORMATION FOR SEQ ID NO: 3:
        (i) SEQUENCE CHARACTERISTICS:
85
             (A) LENGTH: 10 base pairs
86
             (B) TYPE: nucleic acid
87
            (C) STRANDEDNESS: both
88
89
             (D) TOPOLOGY: linear
       (ii) MOLECULE TYPE: cDNA
91
       (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:
                                                                           10
98 ATCGCATGAT
100 (2) INFORMATION FOR SEQ ID NO: 4:
        (i) SEQUENCE CHARACTERISTICS:
102
              (A) LENGTH: 10 base pairs
103
              (B) TYPE: nucleic acid
104
              (C) STRANDEDNESS: both
105
106
              (D) TOPOLOGY: linear
        (ii) MOLECULE TYPE: cDNA
108
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:
113
                                                                             10
115 ATCATGCGAT
117 (2) INFORMATION FOR SEQ ID NO: 5:
         (i) SEQUENCE CHARACTERISTICS:
119
              (A) LENGTH: 65 base pairs
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              (B) TYPE: nucleic acid
121
              (C) STRANDEDNESS: both
122
              (D) TOPOLOGY: linear
123
        (ii) MOLECULE TYPE: cDNA
125
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5:
131
133 AATTCTCGGG ATATCTCAGG ATATCTCAGG ATATCTCAGG ATATCTCGGG
                                                                             60
                                                                             65
135 CTGCA
137 (2) INFORMATION FOR SEQ ID NO: 6:
        (i) SEQUENCE CHARACTERISTICS:
              (A) LENGTH: 43 base pairs
              (B) TYPE: nucleic acid
141
              (C) STRANDEDNESS: single
142
              (D) TOPOLOGY: linear
143
        (ii) MOLECULE TYPE: cDNA
145
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 6:
150
                                                                             43
152 CAACAACAAG AATTCAGGCC TAGCCAGTGC CTCGTCCATT TTT
154 (2) INFORMATION FOR SEQ ID NO: 7:
         (i) SEQUENCE CHARACTERISTICS:
156
157
              (A) LENGTH: 37 base pairs
              (B) TYPE: nucleic acid
158
              (C) STRANDEDNESS: single
159
              (D) TOPOLOGY: linear
160
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37

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162 (ii) MOLECULE TYPE: cDNA

167 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 7:

169 CAACAACAAA AGCTTCTTAC ATGGCCCAGG TGCAGTA

 $W--> 172(-3^{-3})$

file://C:\CRF4\Outhold\VsrI472067A.htm

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/472,067A

DATE: 05/13/2003 TIME: 16:08:28

Input Set : A:\PTO.VSK.txt

Output Set: N:\CRF4\05132003\1472067A.raw

L:6 M:220 C: Keyword misspelled or invalid format, [(i) APPLICANT:]

L:29 M:220 C: Keyword misspelled or invalid format, [(A) APPLICATION NUMBER:]

L:30 M:220 C: Keyword misspelled or invalid format, [(B) FILING DATE:]

L:36 M:238 W: Alpha Fields not Ordered, Reordered [(C) CLASSIFICATION:] of (1)(vi) L:41 M:238 W: Alpha Fields not Ordered, Reordered [(C) CLASSIFICATION:] of (1)(vi)

L:172 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:7